## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

- (i) APPLICANT:
  - (A) NAME: YEDA RESEARCH AND DEVELOPMENT COMPANY LTD.
  - (B) STREET: THE WETZMANN INSTITUTE OF SCIENCE
  - (C) CITY: REHOVOT
  - (E) COUNTRY: ISRAEI
  - (F) POSTAL CODE (ZIP): P.O.BOX 95
  - (G) TELEPHONE: +08 /9470617
  - (H) TELEFAX: +08 9470739
- (ii) TITLE OF INVENTION: ASSAY FOR THE DIAGNOSIS OF SCHIZOPHRENIA EASED ON A NEW PERTIDE
- (iii) NUMBER OF SEQUENCES: 14
  - (iv) COMPUTER READABLE FORM:
    - (A) MEDIUM TYPE: Floppy disk
    - (B) COMPUTER: IBM PC compatible
      - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
      - (D) SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
- (2) INFORMATION FOR SEQ ID NO: 1:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 28 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: unknown
    - (D) TOPOLOGY: unknown
  - (ii) MOLECULE TYPE: peptide
  - (iii) HYPOTHETICAL: NO
    - (iv) ANTI-SENSE: NO
    - (v) FRAGMENT TYPE: internal
    - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
    - Ser Gly Glu Thr Glu Asp Thr Phe Ile Ala Asp Leu Val Val Gly Leu
    - Cys Thr Gly Gln Ile Lys Thr Gly Ala Pro Cys Arg
  - (2) INFORMATION FOR SEQ ID NO: 2:
    - (1) SEQUENCE CHARACTERISTICS:
      - (A) LENGTH: 16 amino acids
      - (3) TYPE: amino acid
      - (C) STRANDEDNESS: unknown
      - (D) TOPOLOGY: wiknown

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- (ii) MOLECULE TYPE: peptide
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- (v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Leu Val Val Gly Leu Cys Thr Cys Gln Ile Lys Thr Gly Pro Ala Cys
1 10 15

(2) INFORMATION FOR SEQ ID NO: 3:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 20 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: unknown
  - (D) TOPOLOGY: unknown
- (ii) MOLECULE TYPE: peptide
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
  - (v) FRAGMENT TYPE: internal
- (xi) SEQUENCE DESCRIPTION: SEP ID NO: 3:

Ile Ala Asp Leu Val Val Gly Leu Cys Thr Gly Gln Ile Lys Thr Gly
1 5 10 15

Ala Pro Cys Arg 20

- (2) INFORMATION FOR SEQ ID NO: 4:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: unknown
    - (D) TOPOLOGY: unknown
  - (ii) MOLECULE TYPE: peptide
  - (111) HYPOTHETICAL: NO
  - (iv) ANTI-SINSE: NO
    - (v) FRAGMENT TYPE: incomal

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
         Ala Asp Leu Val Val Gly Leu Cys Thr Gly Gln Ile Lys Thr Gly Ala
         Pro Cys Arg
    (2)
        INFORMATION FOR SEQ ID NO: 5:
         (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 18 amino acids
              (B) TYPE: amino adid
               (C) STRANDEDNESS: unknown
               (D) TOPOLOGY: unknown
        (ii) MOLECULE TYPE: peptide
(iii) HYPOTHETICAL: NO
        (iv) ANTI-SENSE: NO
         (v) FRAGMENT TYPE: internal
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         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
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         Asp Leu Val Val Gly Leu Cys Thr Gly Gln Ile Lys Thr Gly Ala Pro
O
Cys Arg
     (2) INFORMATION FOR SEQ ID NO: 6:
          (i) SEQUENCE CHARACTERISTICS:
               (A) LENGTH: 17 amino acids
               (B) TYPE: amino acid
               (C) STRANDEDNESS: unknown
               (D) TOPOLOGY: unknown
         (ii) MOLECULE TYPE: peptide
         (iii) EYPOTHETICAL: NO
         (iv) ANTI-SENSE: NO
          (v) FRAGMENT TYPE: internal
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## SUBSTITUTE SHEET (RULE 26)

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Leu Val Val Gly Leu Cys Thr Gly\Gln Ile Lys Thr Gly Ala Bro Cys

(x1) SEQUENCE DESCREPTION: SEQ 4D NO: 6:

Arg

(2) INFORMATION FOR SEQ ID NO: 7:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 17 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS unknown
  - (D) TOPOLOGY: unknown
- (ii) MOLECULE TYPE: pertide
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
  - (v) FRAGMENT TYPE: internal
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

Leu Val Val Gly Leu Cys Thr Gly Gln Ile Lys Thr Gly Pro Ala Cys
1 10 15

Arg

- (2) INFORMATION FOR SEQ ID NO: 8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 17 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: unknown
    - (D) TOPOLOGY: unknown
  - (ii) MOLECULE TYPE: peptide
  - (iii) HYPOTHETICAL: NO
    - (iv) ANTI-SENSE: NO
    - (v), FRAGMENT TYPE: internal
    - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

Leu Val Val Gly Leu Cys Thr Pro Gln Ile Lys Thr Gly Pro Ala Cys
1 5 10 15

Arg

- (2) INFORMATION FOR SEQ ID NO: 9:
  - (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 20 amino acids

- (3) TYPE: amino acid
- (C) STRANDEDNESS: unknown
- (D) TOPCLOGY: unknown
- (ii) MOLECULE TYPE: peptide
- (iii) HYPOTHETICAL: NO
- (iv) ANTI-SENSE: NO
- (v) FRAGMENT TYPE: internal
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

Ser Gly Glu Thr Glu Asp Thr the Ile Ala Asp Leu Val Val Gly Leu

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Cys Thr Gly Gln 20

- (2) INFORMATION FOR SEQ ID NO: 10:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 16 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: unknown
    - (D) TOPOLOGY: unknown
  - (ii) MOLECULE TYPE: peptide
  - (iii) HYPOTHETICAL: NO
  - (iv) ANTI-SENSE: NO
  - (v) FRAGMENT TYPE: internal
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

Val Val Gly Leu Cys Thr Gly Gln Ile Lys Thr Gly Ala Pro Cys Arg
1 10 15

- (2) INFORMATION FOR SEQ ID NO: 11:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 12 amino acids
    - (5) TYPE: amino acid
    - (C) STRANDEDNESS: unknown
    - (D) TOPOLOGY: unknown
  - (ii) MOLECULE TYPE: peptide
  - (iii) HYPOTHETICAL: NO
  - (iv) ANTI-SENSE: NO

(v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

Cys Thr Gly Gln Ile Lys Thr Gly Ala Pro Cys Arg

- (2) INFORMATION FOR SEQ ID NO: 12:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 16 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: unknown
    - (D) TOPOLOGY: unknown
  - (ii) MOLECULE TYPE: peptide
  - (iii) HYPOTHETICAL: NO
  - (iv) ANTI-SENSE: NO
  - (v) FRAGMENT TYPE: internal
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

Leu Val Val Gly Leu Cys Thr Gly Gln Ile Lys Thr Gly Ala Pro Cys
1 10 15

- (2) INFORMATION FOR SEQ ID NO: 13:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 15 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: unknown
    - (D) TOPOLOGY: unknown
  - (ii) MOLECULE TYPE: peptide
  - (iii) HYPOTHETICAL: NO
  - (iv) ANTI-SENSE: NO
    - (v) FRAGMENT TYPE: internal
  - (ki) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

Leu Val Val Gly Leu Cys Thr/Gly Gln Ile Lys Thr Gly Ala Pro 1 10 13

(2) INFORMATION FOR SEQ ID NO: 14

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 16 amino acids
    (B) TYPE amino acid

  - (C) STRANDEDNESS: unknown
  - (D) TOPOLOGY: unknown
- (ii) MOLECULE TYPE: peptide
- (iii) HYPOTHETICAL:\NO
- (iv) ANTI-SENSE: NO
- (v) FRAGMENT TYPE: internal

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

Leu Val Val Gly Leu Cys Thr Gly Gln Ile Lys Thr Gly Pro Ala Cys

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